

# SCORE Search Results Details for Application 10501412 and Search Result 20070411\_152412\_us-10-501-412- 1\_copy\_1\_445.rng.

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This page gives you Search Results detail for the Application 10501412 and Search Result 20070411\_152412\_us-10-501-412-1\_copy\_1\_445.rng.

[Go Back to previous page](#)

GenCore version 6.2  
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OM nucleic - nucleic search, using sw model

Run on: April 12, 2007, 07:02:58 ; Search time 313 Seconds  
(without alignments)  
10536.787 Million cell updates/sec

Title: US-10-501-412-1\_COPY\_1\_445  
Perfect score: 445  
Sequence: 1 accacaaacacuccaguug.....agucucguagaccguagcac 445

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 5620219 seqs, 3705283702 residues

Total number of hits satisfying chosen parameters: 11240438

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : N\_Geneseq\_200701:\*  
1: geneseqn1980s:\*  
2: geneseqn1990s:\*  
3: geneseqn2000s:\*  
4: geneseqn2001as:\*  
5: geneseqn2001bs:\*  
6: geneseqn2002as:\*  
7: geneseqn2002bs:\*  
8: geneseqn2003as:\*  
9: geneseqn2003bs:\*  
10: geneseqn2003cs:\*  
11: geneseqn2003ds:\*  
12: geneseqn2004as:\*  
13: geneseqn2004bs:\*  
14: geneseqn2005s:\*  
15: geneseqn2006s:\*  
16: geneseqn2007s:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.